

SEQUENCE LISTING

☐

☐

<110> Imperial Cancer Research Technology Limited

☐

☐

<120> Polypeptides and their use in therapy

☐

☐

<130> IMPW/P18999PC

☐

☐

<140>

☐

<141>

☐

☐

<160> 2

☐

☐

<170> PatentIn Ver. 2.0

☐

☐

<210> 1

☐

<211> 49

☐

<212> DNA

☐

<213> Artificial Sequence

☐

☐

<220>

☐

<223> Description of Artificial Sequence:double stranded

☐

oligonucleotide

☐

☐

<400> 1

☐

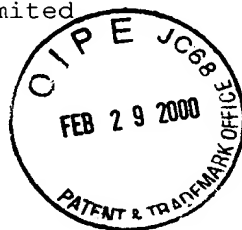
cacagtcagg acatcatcat catcatcatt aaggatcctc tagaggtag

☐

☐

<210> 2

☐



<211> 728

□

<212> PRT

□

<213> h. sapiens

□

□

<400> 2

□

Met Trp Val Thr Lys Leu Leu Pro Ala Leu Leu Leu Gln His Val Leu

□

1

5

10

15

□

□

Leu His Leu Leu Leu Leu Pro Ile Ala Ile Pro Tyr Ala Glu Gly Gln

□

20

25

30

□

□

Arg Lys Arg Arg Asn Thr Ile His Glu Phe Lys Lys Ser Ala Lys Thr

□

35

40

45

□

□

Thr Leu Ile Lys Ile Asp Pro Ala Leu Lys Ile Lys Thr Lys Lys Val

□

50

55

60

□

□

Asn Thr Ala Asp Gln Cys Ala Asn Arg Cys Thr Arg Asn Lys Gly Leu

□

65

70

75

80

□

□

Pro Phe Thr Cys Lys Ala Phe Val Phe Asp Lys Ala Arg Lys Gln Cys

□

85

90

95

□

□

Leu Trp Phe Pro Phe Asn Ser Met Ser Ser Gly Val Lys Lys Glu Phe

□

100

105

110

□

□

Gly His Glu Phe Asp Leu Tyr Glu Asn Lys Asp Tyr Ile Arg Asn Cys

□

115

120

125

□

☐ Ile Ile Gly Lys Gly Arg Ser Tyr Lys Gly Thr Val Ser Ile Thr Lys

☐ 130 135 140

☐
☐ Ser Gly Ile Lys Cys Gln Pro Trp Ser Ser Met Ile Pro His Glu His

☐ 145 150 155 160

☐
☐ Ser Phe Leu Pro Ser Ser Tyr Arg Gly Lys Asp Leu Gln Glu Asn Tyr

☐ 165 170 175

☐
☐ Cys Arg Asn Pro Arg Gly Glu Glu Gly Gly Pro Trp Cys Phe Thr Ser

☐ 180 185 190

☐
☐ Asn Pro Glu Val Arg Tyr Glu Val Cys Asp Ile Pro Gln Cys Ser Glu

☐ 195 200 205

☐
☐ Val Glu Cys Met Thr Cys Asn Gly Glu Ser Tyr Arg Gly Leu Met Asp

☐ 210 215 220

☐
☐ His Thr Glu Ser Gly Lys Ile Cys Gln Arg Trp Asp His Gln Thr Pro

☐ 225 230 235 240

☐
☐ His Arg His Lys Phe Leu Pro Glu Arg Tyr Pro Asp Lys Gly Phe Asp

☐ 245 250 255

☐
☐ Asp Asn Tyr Cys Arg Asn Pro Asp Gly Gln Pro Arg Pro Trp Cys Tyr

☐ 260 265 270

☐

☐

Thr Leu Asp Pro His Thr Arg Trp Glu Tyr Cys Ala Ile Lys Thr Cys

□

275

280

285

□

□

Ala Asp Asn Thr Met Asn Asp Thr Asp Val Pro Leu Glu Thr Thr Glu

□

290

295

300

□

□

Cys Ile Gln Gly Gln Gly Glu Gly Tyr Arg Gly Thr Val Asn Thr Ile

□

305

310

315

320

□

□

Trp Asn Gly Ile Pro Cys Gln Arg Trp Asp Ser Gln Tyr Pro His Glu

□

325

330

335

□

□

His Asp Met Thr Pro Glu Asn Phe Lys Cys Lys Asp Leu Arg Glu Asn

□

340

345

350

□

□

Tyr Cys Arg Asn Pro Asp Gly Ser Glu Ser Pro Trp Cys Phe Thr Thr

□

355

360

365

□

□

Asp Pro Asn Ile Arg Val Gly Tyr Cys Ser Gln Ile Pro Asn Cys Asp

□

370

375

380

□

□

Met Ser His Gly Gln Asp Cys Tyr Arg Gly Asn Gly Lys Asn Tyr Met

□

385

390

395

400

□

□

Gly Asn Leu Ser Gln Thr Arg Ser Gly Leu Thr Cys Ser Met Trp Asp

□

405

410

415

□

□

Lys Asn Met Glu Asp Leu His Arg His Ile Phe Trp Glu Pro Asp Ala

□

	420	425	430
□			
□			
Ser	Lys	Leu	Asn
□	Glu	Asn	Tyr
	Cys	Arg	Asn
	Pro	Asp	Asp
	Asp	Ala	His
	435	440	445
□			
□			
Gly	Pro	Trp	Cys
□	Tyr	Thr	Gly
	Asn	Pro	Leu
	Ile	Pro	Trp
	Asp	Tyr	Cys
	450	455	460
□			
□			
Pro	Ile	Ser	Arg
□	Cys	Glu	Gly
	Asp	Thr	Thr
	Pro	Thr	Ile
	Val	Asn	Leu
465	470	475	480
□			
□			
Asp	His	Pro	Val
□	Ile	Ser	Cys
	Ala	Lys	Thr
	Lys	Gln	Leu
	Arg	Val	Val
	485	490	495
□			
□			
Asn	Gly	Ile	Pro
□	Thr	Arg	Thr
	Asn	Ile	Gly
	Trp	Met	Val
	Ser	Leu	Arg
	500	505	510
□			
□			
Tyr	Arg	Asn	Lys
□	His	Ile	Cys
	Gly	Gly	Ser
	Leu	Ile	Lys
	Glu	Ser	Trp
	515	520	525
□			
□			
Val	Leu	Thr	Ala
□	Arg	Gln	Cys
	Phe	Pro	Ser
	Arg	Asp	Leu
	Lys	Asp	Tyr
	530	535	540
□			
□			
Glu	Ala	Trp	Leu
□	Gly	Ile	His
	Asp	Val	His
	Gly	Arg	Gly
	Asp	Glu	Lys
545	550	555	560
□			
□			
Cys	Lys	Gln	Val
□	Leu	Asn	Val
	Ser	Gln	Leu
	Val	Tyr	Gly
	Pro	Glu	Gly
	565	570	575
□			

☐ Ser Asp Leu Val Leu Met Lys Leu Ala Arg Pro Ala Val Leu Asp Asp

☐ 580 585 590

☐ Phe Val Ser Thr Ile Asp Leu Pro Asn Tyr Gly Cys Thr Ile Pro Glu

☐ 595 600 605

☐ Lys Thr Ser Cys Ser Val Tyr Gly Trp Gly Tyr Thr Gly Leu Ile Asn

☐ 610 615 620

☐ Tyr Asp Gly Leu Leu Arg Val Ala His Leu Tyr Ile Met Gly Asn Glu

☐ 625 630 635 640

☐ Lys Cys Ser Gln His His Arg Gly Lys Val Thr Leu Asn Glu Ser Glu

☐ 645 650 655

☐ Ile Cys Ala Gly Ala Glu Lys Ile Gly Ser Gly Pro Cys Glu Gly Asp

☐ 660 665 670

☐ Tyr Gly Gly Pro Leu Val Cys Glu Gln His Lys Met Arg Met Val Leu

☐ 675 680 685

☐ Gly Val Ile Val Pro Gly Arg Gly Cys Ala Ile Pro Asn Arg Pro Gly

☐ 690 695 700

☐ Ile Phe Val Arg Val Ala Tyr Tyr Ala Lys Trp Ile His Lys Ile Ile

☐ 705 710 715 720

☐

☐

Leu Thr Tyr Lys Val Pro Gln Ser

☐

725

☐

☐

☐

☐

Sub
B3
cont.
